OIPE

CRF Errors Corrected by the STIC Systems Branch

CRF Processing Date: 6/9/2003

Serial Number: 09/939,853A

Edited by:

Verified by: (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING DATE: 06/09/2003
PATENT APPLICATION: US/09/939,853A TIME: 13:56:26

Input Set : N:\AMC\SEQUENCE LISTING CURA-399 US.DOC
Output Set: N:\CRF4\06092003\I939853A.raw

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 3 <110> APPLICANT: Burgess et al.
 5 <120> TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
 7 <130> FILE REFERENCE: CURA-399
 9 <140> CURRENT APPLICATION NUMBER: 09/939,853A
10 <141> CURRENT FILING DATE: 2001-08-27
12 <150> PRIOR APPLICATION NUMBER: 60/228,191
13 <151> PRIOR FILING DATE: 2000-08-25
15 <150> PRIOR APPLICATION NUMBER: 60/267,300
16 <151> PRIOR FILING DATE: 2001-02-08
18 <150> PRIOR APPLICATION NUMBER: 60/269,961
19 <151> PRIOR FILING DATE: 2001-02-20
21 <150> PRIOR APPLICATION NUMBER: 60/277,337
22 <151> PRIOR FILING DATE: 2001-03-20
24 <160> NUMBER OF SEQ ID NOS: 150
26 <170> SOFTWARE: PatentIn Ver. 2.1
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 1065
30 <212> TYPE: DNA
31 <213> ORGANISM: Homo sapiens
33 <400> SEQUENCE: 1
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35 tcgatggagt gtctgtgcag ttgggagcag aggctctgcc tctctggtcc taccttggg 120
36 agtgaaatag tgaccgtccc ttgagcccg agagctgggg agaaggccgt gcctgttaac 180
37 agctgcctgg acctctctg cagacgcagca gagagaggcg gggctggagg agatgttgcc 240
38 aagaacctaa ggctgaaagt catgcttgcc atctgttcca ccttggctcg agatacttca 300
39 gaatccagct ctttcttaaa ggtggacccc cggaagaagc agatcacctt gtacgatccc 360
40 ctgacttgtg caggtcaaaa tgccttccaa aagagaggca accaggttcc tccaaagatg 420
41 tttgccttcg atccagtttt tccacaagac gcttctcagg ctgaagtgtg tgcaggcacc 480
42 ttggcagagg tgatccagtc tgtgtccaac ggggcagatg gctgcgtgtt ctgtttcggc 540
43 cacgccaaac tgcgaaaatc ctaccaccat gatcggaaagg atgattccat gcagaacctg 600
44 tccatcattc ctgtgccat ctctggctc ttcaagctca taaacgaacg caaggaaaag 660
45 accgcggcca gtttctcagt ccggctttcc gccgtggaag tgtgggggaa ggaggagaac 720
46 ctgcaggacc tgctgtcgga gctgcccacg ggcagcctgc aggacggcca gtccccgggc 780
47 ctgtcctct gtcaggacc catcgcggc acgcagctgc agaaccagag cgagctgcgg 840
48 gccccaccg cagaggaggc tgccttttc ctggatgccg ccattgcctc ccgcaggagc 900
49 taccaacagg actgtgatga ggacgaccac cgcaactcac acgtgttctt cacactgcac 960
50 atctgccact acggatgga ggagagcggg aaagggggaa ttctgctttc gatttggaat 1020
51 ctgaaagtag ggggaaatct tgaaaacaag gaaacagttc attaa 1065
54 <210> SEQ ID NO: 2
55 <211> LENGTH: 354
56 <212> TYPE: PRT
57 <213> ORGANISM: Homo sapiens
59 <400> SEQUENCE: 2
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RAW SEQUENCE LISTING DATE: 06/09/2003
PATENT APPLICATION: US/09/939,853A TIME: 13:56:26

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Output Set: N:\CRF4\06092003\I939853A.raw

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60 Met Thr Gly Leu Leu Leu Leu Ser Leu Gln Ser Gly Cys Val Ala Ala
61  1               5                  10                  15
62 Ile Thr Ser Met Ser Met Glu Cys Leu Cys Ser Leu Gly Ala Arg Leu
63              20                  25                  30
64 Cys Leu Ser Arg Ser Thr Leu Gly Ser Glu Ile Val Thr Val Pro Leu
65          35                  40                  45
66 Ser Pro Arg Ala Gly Glu Lys Ala Val Pro Val Asn Ser Cys Leu Asp
67      50                  55                  60
68 Pro Leu Trp Arg Ala Ala Glu Arg Gly Gly Ala Gly Gly Asp Val Ala
69  65                  70                  75                  80
70 Lys Asn Leu Arg Val Lys Val Met Leu Arg Ile Cys Ser Thr Leu Ala
71                  85                  90                  95
72 Arg Asp Thr Ser Glu Ser Ser Ser Phe Leu Lys Val Asp Pro Arg Lys
73              100                 105                 110
74 Lys Gln Ile Thr Leu Tyr Asp Pro Leu Thr Cys Glu Gly Gln Asn Ala
75          115                 120                 125
76 Pro Gln Lys Arg Gly Asn Gln Val Pro Pro Lys Met Pro Ala Phe Asp
77      130                 135                 140
78 Ala Val Phe Pro Gln Asp Ala Ser Gln Ala Glu Val Cys Ala Gly Thr
79  145                 150                 155                 160
80 Val Ala Glu Val Ile Gln Ser Val Val Asn Gly Ala Asp Gly Cys Val
81                  165                 170                 175
82 Phe Cys Phe Gly His Ala Lys Leu Gly Lys Ser Tyr Thr Met Ile Gly
83              180                 185                 190
84 Lys Asp Asp Ser Met Gln Asn Leu Gly Ile Ile Pro Cys Ala Ile Ser
85          195                 200                 205
86 Trp Leu Phe Lys Leu Ile Asn Glu Arg Lys Glu Lys Thr Gly Ala Arg
87      210                 215                 220
88 Phe Ser Val Arg Val Ser Ala Val Glu Val Trp Gly Lys Glu Glu Asn
89  225                 230                 235                 240
90 Leu Arg Asp Leu Leu Ser Glu Val Ala Thr Gly Ser Leu Gln Asp Gly
91                  245                 250                 255
92 Gln Ser Pro Gly Val Tyr Leu Cys Glu Asp Pro Ile Cys Gly Thr Gln
93              260                 265                 270
94 Leu Gln Asn Gln Ser Glu Leu Arg Ala Pro Thr Ala Glu Lys Ala Ala
95          275                 280                 285
96 Phe Phe Leu Asp Ala Ala Ile Ala Ser Arg Arg Ser His Gln Gln Asp
97      290                 295                 300
98 Cys Asp Glu Asp Asp His Arg Asn Ser His Val Phe Phe Thr Leu His
99  305                 310                 315                 320
100 Ile Tyr Gln Tyr Arg Met Glu Lys Ser Gly Lys Gly Gly Ile Leu Leu
101                 325                 330                 335
102 Ser Ile Trp Asn Leu Lys Val Gly Arg Asn Leu Glu Asn Lys Glu Thr
103             340                 345                 350
104 Val His
105 <210> SEQ ID NO: 3
106 <211> LENGTH: 366
107 <212> TYPE: PRT
108 <213> ORGANISM: Homo sapiens
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RAW SEQUENCE LISTING DATE: 06/09/2003
PATENT APPLICATION: **US/09/939,853A** TIME: 14:56:26

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Output Set: **N:\CRF4\06092003\I939853A.raw**

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137   1               5                  10                  15
138 Pro Arg Arg Ala Ala Thr Ala Ala Val Pro Lys Met Phe Ala Phe Asp
139             20                  25                  30
140 Ala Val Phe Pro Gln Asp Ser Glu Gln Ala Glu Val Cys Ser Gly Thr
141         35                  40                  45
142 Val Ala Asp Val Leu Gln Ser Val Val Ser Gly Ala Asp Gly Cys Ile
143     50                  55                  60
144 Phe Ser Phe Gly His Met Ser Leu Gly Lys Ser Tyr Thr Met Ile Gly
145 65                  70                  75                  80
146 Lys Asp Ser Ser Pro Gln Ser Leu Gly Ile Val Pro Cys Ala Ile Ser
147                 85                  90                  95
148 Trp Leu Phe Arg Leu Ile Glu Glu Arg Arg Glu Arg Thr Gly Thr Arg
149             100                 105                 110
150 Phe Ser Val Arg Val Ser Ala Val Glu Val Cys Gly Arg Asp Gln Ser
151         115                 120                 125
152 Leu Arg Asp Leu Leu Ala Glu Val Ala Pro Gly Ser Leu Gln Asp Thr
153     130                 135                 140
154 Gln Ser Pro Gly Val Tyr Leu Arg Glu Asp Pro Val Cys Gly Ala Gln
155 145                 150                 155                 160
156 Leu Gln Asn Gln Ser Glu Leu Arg Ala Pro Thr Ala Glu Lys Ala Ala
157                 165                 170                 175
158 Phe Tyr Leu Asp Ala Ala Leu Ala Ala Arg Ser Thr Ser Arg Ala Gly
159             180                 185                 190
160 Cys Gly Glu Asp Ala Arg Arg Ser Ser His Met Leu Phe Thr Leu His
161         195                 200                 205
162 Val Tyr Gln Tyr Arg Met Glu Lys Cys Gly Arg Gly Gly Met Ser Gly
163     210                 215                 220
164 Gly Arg Ser Arg Leu His Leu Ile Asp Leu Gly Ser Cys Glu Ala Ala
165 225                 230                 235                 240
166 Ala Gly Arg Ala Gly Glu Ala Ala Gly Gly Pro Leu Cys Leu Ser Leu
167                 245                 250                 255
168 Ser Ala Leu Gly Ser Val Ile Leu Ala Leu Val Asn Gly Ala Lys His
169             260                 265                 270
170 Val Pro Tyr Arg Asp His Arg Leu Thr Met Leu Leu Arg Glu Ser Leu
171         275                 280                 285
172 Ala Thr Ala Gly Cys Arg Thr Thr Met Ile Ala His Val Ser Asp Ala
173     290                 295                 300
174 Pro Ala Gln His Ala Glu Thr Leu Ser Thr Val Gln Leu Ala Ala Arg
175 305                 310                 315                 320
176 Ile His Arg Leu Arg Arg Lys Lys Ala Lys Tyr Ala Ser Ser Ser Ser
177                 325                 330                 335
178 Gly Gly Glu Ser Ser Cys Glu Glu Gly Arg Ala Arg Arg Pro Pro His
179             340                 345                 350
180 Leu Arg Pro Phe His Pro Arg Thr Val Ala Leu Asp Pro Asp
181         355                 360                 365
182 <210> SEQ ID NO: 4
183 <211> LENGTH: 130
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RAW SEQUENCE LISTING DATE: 06/09/2003
PATENT APPLICATION: US/09/939,853A TIME: 18:06:26

Input Set : N:\AMC\SEQUENCE LISTING CURA-399 US.DOC
Output Set: N:\CRF4\06092003\I939853A.raw

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108 <212> TYPE: PRT
109 <213> ORGANISM: Mus musculus
111 <400> SEQUENCE: 4
112 Pro Ala Pro Thr Gly Lys Ser Tyr Thr Met Ile Gly Arg Asp Asp Ser
113   1               5                  10                  15
115 Met Gln Asn Leu Gly Ile Ile Pro Cys Ala Ile Ser Trp Leu Phe Lys
116             20                  25                  30
118 Leu Ile Asn Glu Arg Lys Glu Lys Thr Gly Ala Arg Phe Ser Val Arg
119         35                  40                  45
121 Ile Ser Ala Val Glu Val Trp Gly Lys Glu Glu Asn Leu Arg Asp Leu
122     50                  55                  60
124 Leu Ser Glu Val Ala Thr Gly Ser Leu Gln Asp Gly Gln Ser Pro Gly
125 65                  70                  75                  80
127 Val Tyr Leu Cys Glu Asp Pro Ala Glu Lys Ala Ala Phe Phe Leu Asp
128                 85                  90                  95
130 Ala Ala Ile Asn Ser Arg Arg Ser Asn Gln Gln Asp Cys Asp Glu Asp
131             100                 105                 110
133 Asp His Arg Asn Ser His Met Leu Phe Thr Leu His Ile Tyr Gln Tyr
134         115                 120                 125
136 Arg Met
137     130
140 <210> SEQ ID NO: 5
141 <211> LENGTH: 147
142 <212> TYPE: PRT
143 <213> ORGANISM: Mus musculus
145 <400> SEQUENCE: 5
146 Gly Lys Ser Tyr Thr Met Ile Gly Lys Asp Ser Ser Pro Gln Ser Leu
147   1               5                  10                  15
149 Gly Ile Val Pro Cys Ala Ile Ser Trp Leu Phe Arg Leu Ile Asp Glu
150             20                  25                  30
152 Arg Lys Glu Arg Leu Gly Thr Arg Phe Ser Ile Arg Val Ser Ala Val
153         35                  40                  45
155 Glu Val Cys Gly His Asp Gln Ser Leu Arg Asp Leu Leu Ala Glu Val
156     50                  55                  60
158 Ala Ser Gly Ser Leu Gln Asp Thr Gln Ser Pro Gly Val Tyr Leu Arg
159 65                  70                  75                  80
161 Glu Asp Pro Val Cys Gly Thr Gln Leu Gln Asn Gln Asn Glu Leu Arg
162                 85                  90                  95
164 Ala Pro Thr Ala Glu Lys Ala Ala Phe Tyr Leu Asp Ala Ala Leu Ala
165             100                 105                 110
167 Ala Arg Ser Thr Ser Arg Ala Gly Cys Gly Glu Glu Ala Arg Arg Ser
168         115                 120                 125
170 Ser His Met Leu Phe Thr Leu His Val Tyr Gln Tyr Arg Val Glu Lys
171     130                 135                 140
173 Cys Gly Gln
174 145
177 <210> SEQ ID NO: 6
178 <211> LENGTH: 302
179 <212> TYPE: PRT
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RAW SEQUENCE LISTING DATE: 06/09/2003
PATENT APPLICATION: US/09/939,853A TIME: 13:56:26

Input Set : N:\AMC\SEQUENCE LISTING CURA-399 US.DOC
Output Set: N:\CRF4\06092003\I939853A.raw

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280 <213> ORGANISM: Drosophila melanogaster
282 <400> SEQUENCE: 6
283 Met Ala Thr Thr Ser Thr Ser Asn Met Ser Arg Asn Gly Gly Phe Cys
284   1               5                  10                  15
286 Gly Ala Leu Gln Arg Ala Pro Pro Pro Met Pro Pro Thr Leu Ile Arg
287             20                  25                  30
289 Arg Leu Ser Ser Arg Glu Cys Tyr Gly Val Gly Lys Val Lys Val Met
290         35                  40                  45
292 Leu Arg Val Ala Asp Arg Asp Arg Asn Ser Gly Gly Thr Glu Pro Asp
293     50                  55                  60
295 Phe Met Ala Leu Asp Lys Lys Lys Arg Gln Val Thr Leu Thr Asp Pro
296 65                  70                  75                  80
298 Arg Thr Ala Cys Pro Pro Pro Gln Ala Ala Gln Glu Arg Ala Pro Met
299                 85                  90                  95
301 Val Ala Ala Pro Lys Met Phe Ala Phe Asp Asn Leu Phe Thr Gly Glu
302             100                 105                 110
304 Asp Lys Gln Ser Asp Val Cys Ala Ser Ala Leu Ser Glu Val Ile Pro
305         115                 120                 125
307 Ala Val Leu Glu Gly Ser Asp Gly Cys Leu Leu Ala Met Gly Tyr Pro
308     130                 135                 140
310 Ala Thr Gly Gln Ala Gln Thr Val Leu Gly Glu Leu Gly Gly Gly Ser
311 145                 150                 155                 160
313 Gly Ser Gly Ser Ala Ser Gly Ser Gly Val Ala Cys Ser Leu Gly Ala
314                 165                 170                 175
316 Ala Pro Cys Ala Ile Ala Trp Leu Tyr Lys Gly Ile Gln Glu Arg Arg
317             180                 185                 190
319 Gln Lys Ser Gly Ala Arg Phe Ser Val Arg Val Ser Ala Val Gly Val
320         195                 200                 205
322 Ser Ala Thr Lys Pro Asp Ala Leu Ser Gln Asp Leu Leu Ile Ser His
323     210                 215                 220
325 Ala Ala Glu Tyr Gly Val Tyr Ser His Ile Lys Pro Asn Ala Leu Phe
326 225                 230                 235                 240
328 Ile His Ser Pro Leu Leu Phe Phe Trp Ser Gln Tyr Trp Asn Ser Gly
329                 245                 250                 255
331 Ser Asp Tyr Gly Tyr Thr Glu Ser Asp Asp Ser Pro Gly Ile Tyr Leu
332             260                 265                 270
334 Arg Asp Asp Phe Leu Ala Val Gln Arg Asn Tyr Val His Pro Pro Pro
335         275                 280                 285
337 Ser Val Arg Pro Phe Ser Ser Thr Gln Arg Ser Pro Asp Ala
338     290                 295                 300
341 <210> SEQ ID NO: 7
342 <211> LENGTH: 359
343 <212> TYPE: PRT
344 <213> ORGANISM: Caenorhabditis elegans
346 <400> SEQUENCE: 7
347 Met Glu Ala Cys Ser Ser Lys Thr Ser Leu Leu Leu His Ser Pro Leu
348   1               5                  10                  15
350 Arg Thr Ile Pro Lys Leu Arg Leu Cys Ala Ser Ile Ser Ser Glu Asp
351             20                  25                  30
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VERIFICATION SUMMARY DATE: 06/09/2003
PATENT APPLICATION: US/09/939,853A TIME: 18:56:27

Input Set : N:\AMC\SEQUENCE LISTING CURA-399 US.DOC
Output Set: N:\CRF4\06092003\I939853A.raw